



## SEQUENCE LISTING

4110 Scholler, Nathalie B.  
4111 Osis, Mary L.  
4112 Hellstrom, Ingegerd  
4113 Hellstrom, Karl Erik

4114 SURFACE RECEPTOR ANTIGEN VACCINES

4130 230033.409

4140 US 09/441,411  
4141 1999-11-16

4160 16

4170 FastSEQ for Windows Version 4.0

4210 1  
4211 19  
4212 LNA  
4213 Artificial Sequence

4220  
4221 PCR primer

4400 1  
gtatgcttat ggcttgcaat tgtcagttg

29

4210 2  
4211 19  
4212 LNA  
4213 Artificial Sequence

4220  
4221 PCR primer

4400 1  
gtatgcatct aaaggaagac ggtctgttc

29

4210 3  
4211 19  
4212 LNA  
4213 Artificial Sequence

4220  
4221 PCR primer

4400 3  
ggaagcttgt tccagaactt acggaag

27

4210 4

RECEIVED  
SEP 24 2002  
TECH CENTER 1600/2900

AC11: 26  
 AC12: DNA  
 AC13: Artificial Sequence

AC20:  
 AC23: PCR primer

AC40: 4  
 agatgatat ttcctcaggg tctcac

26

AC10: 5  
 AC11: 4473  
 AC12: DNA  
 AC13: Homo sapiens

AC40: 5  
 aaggaggagt aacctggggt cctttgggtg gggcccgggg cagccggggg ccccttccca 60  
 gggggccttt tactgagcgg cgggcccggc ccccaacctt cgcagcaacc cgggcccggc 120  
 gcccctccag cggggtccag cgggagccat ggggcccggg cggcagttag caccatggag 180  
 ctgggggggt tgtgcccgtg ggggtctctc ctggcctctt tggccccggg agccgggagg 240  
 aaccaagtgt gcacgggcac agacatgaag ctggggctcc ctggccagtcc cggagccca 300  
 ctggacatgc tccggccact ctaccagggc tggcagggtg tgcagggaaa cctggaaact 360  
 aactaactgc ccaccaatgc cagctctgtc ttcttgacgg atatccaggg ggtgcagggg 420  
 taagtgtcca tggctcacaa ccaagtggag cagggtccca tgcagagggt gggattgtg 480  
 cgagjcaacc agctcttga ggaacaactat gcccggggc tctagagaaa tggagacccg 540  
 ctgaacata ccacccctgt cacagggggc tcccaggag gctggggga gctgcagctt 600  
 cgaagctcca cagagatctt gaaaggaggg gtcttgatcc aggggaacc ccagctctg 660  
 taccaggaca cgtttttgtt gaaggacata tcccaacaga acaaccagct ggtctcaca 720  
 ctgatagaca ccaaccgctc tggggcctgc caccctctgt ctccgatgtt caagggtcc 780  
 ctgtctgtgg gagaggtctt tgaagattgt cagagctga cggccactgt ctgtgcccgt 840  
 ggtctgtgccc gctgcagggt ggcactggcc actgaactgt ggcctgagca gtgtgctg 900  
 ggtgcacgg gcccacaagg ctctgaactg ctggcctgcc tccacttcaa ccacagtgg 960  
 atctgtgagc tgcactggcc agccctggct accataaca cagacaagtt tgaatccatg 1020  
 ccccaatccc agggccgggtt tacattcggc ggcagctgtg tgaactgctt tccctacaac 1080  
 tacctttcta cggacgtggg atcttgacc ctggtctgcc ccttgcaaaa ccaagaggtg 1140  
 acagcagagg atggaacaca ggggtgtgag aagtgcagca agccctgtgc ccagctgtg 1200  
 tatgttctgg gcatggagca ctggcagag gtgagggcag ttaaccagtgc caatatccag 1260  
 gagtctgtgt gctgcaggaa gatctttggg agcctggcat ttctggcggg gagctttgat 1320  
 ggggacccag cctcccaacc tggcccgttc cagccagagg agctccaagt gtttgagact 1380  
 ctgggaagaga tccagggtta cctatacact tccagatggc cggacagcct gctgacctc 1440  
 agcgtcttcc agaacctgca agtaatccgg ggacgaatto tgcacaatgg cgcctactcg 1500  
 ctgaacctgc aagggctggg catcagctgg ctggggctgc gctcaactgag ggaactgggg 1560  
 agtgtaactg cctccatcca ccataacacc cactctctgt tcttgcaacc ggtgcctgg 1620  
 gaccagctct ttcggaaacc gcaccaagct ctgctccaca ctggcaaccg gccagaggac 1680  
 gagtgtgtgg ggcagggggt ggcctgcac cagctgtggt cccgagggca ctgctgggg 1740  
 ccagggccca cctagtgtgt caactgcagg cagttcttct ggggcccagg gtggtggag 1800  
 gaatggcgag tactgcagg gctcccagg gagtacttga atggcaggca ctgtttggcg 1860  
 tggcaccctg agtgcaggcc ccagaatggc tcaagtgaact gtttgggacc ggaggtctgac 1920  
 cagtgtgtgg cctgtgcaca ctataaggac cctccctctt ggcggggccg ctgcccagc 1980  
 ggtgtgaaac ctgaactctc ctacatgccc atctgggaag ttccagatga ggagggggca 2040  
 tggcagcctt gcccacatca ctgcacccac tccctgtgtg aactggatga caagggtctg 2100  
 ccggccagag agagagccag cctcttgacg tccatcatct ctgggtgtgt tggcattctg 2160  
 ctgggtcgtg tcttgggggg ggtctttggg atctcatca agcgaaggca gcagaagatc 2220  
 cgggaagtaca cgtgctggag actgctgcag gaaacggagc tgggtggagcc gctgacacct 2280

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agcggagcga tgcacaacca ggcgcagatg cggatcctga aagagacgga gctgaggaag 2340
gtgaagggtg ttggatcttg cgtttctggc acagtctaca agggcatctg gatccctgat 2400
ggggagaatg tgaataattcc agtggccatc aaagtgtga gggaaaacac atcccccaaa 2460
gcaacaaaag aaattcttaga cgaagcctac gtgatggctg gtgtgggctc cccatattgtc 2520
ccccgcttc tgggcatctg cctgacatcc acggtgcagc tgggtgacaca gcttatgccc 2580
tatggctgoc tortagacca tgtccgggaa aacccgggac gctgggctc ccaggacctg 2640
ctgaactggt gtatgcagat tgcacagggt atgagctacc tggaggatgt gggctcgtta 2700
cacagggaat tggccgctcg gaactgtctg gtcaagagtc ccaacctatgt caaaattaca 2760
gaactcgggg tggctcggct gctggacatt gacgagacag agtacctatgc agatgggggc 2820
aaggtgcaca tcaagtggat ggcgtggag tccattctcc gcggcggtt caccacacag 2880
agtgatgtgt ggagttatgg tgtgaactgt tgggagctga tgacttttgg ggcacacac 2940
taagatggga tcccagcccg ggagatccct gacctgttg aaaaggggga ggggtgccc 3000
cagcccccca tctgcaacct tgatgtctac atgacatgg tcaaatgttg gatgattgac 3060
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ccccagcgtt ttgttgctac ccagaatgag gacttggcc cagccagtcc cttggacagc 3180
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tatctggtac ccagcaggg cttctctctgt ccagaacctg ccccggggcg tgggggcatg 3300
gtccacacaa ggcacccgag ctcatctacc aggagtggg gtggggacct gacactaggg 3360
ctgagacccc ctgaagagga ggcacccagg tctccactgg caccctccga aggggtggtg 3420
tccgatgtat ttgatgggga cctgggaatg ggggagacca aggggtgga aagcctcccc 3480
acacatgacc ccagccctct acagcgggac agtgaggacc ccacagtacc cctgcctct 3540
gagactgatg gtaacttgc cccctgacc tgpagcccc agcctgaata tgtgaacag 3600
ccagatgttc ggccccagcc cctctcgccc cgagaggcc cctgctctg tgcctgacct 3660
gactgttcca ctctggaaa gcccagaact ctctccccag ggaagaatgg ggtcgtcaaa 3720
gaactttttg cctctggggg tgcctggag aaaccccgagt acctgacac ccaggaggga 3780
gctgcctctc agcccaaccc tctctctgcc ttcaggccag ccttcgacaa cctctattac 3840
tgggacccag accccacaga gcggggggct ccacccagca cctcaaaag gacacctacg 3900
gcacagaaac cagagtacct ggtctggac gtgcctgtgt gaaccagaag gccaagtccg 3960
cagaagccct gatgtgtct cagggagcag ggaagcctg acctctgtgt gcatcaagag 4020
gtgggagggc cctccagacc cctccagggg aacctgcbat gccaggaccc tgtctaaagg 4080
aaccttccct cctgcttgag tccccagatg gctgggaagg gtccagcctc gttggagag 4140
gaacagcact ggggagctct tctggactct gaggccctgc ccaatgagac tctagggtcc 4200
agtggatgoc acagcccagc ttggcccttt ccttcagat cctgggtact gaaagcctta 4260
gggaagctgg cctgagaggg gaaagggccc taaggagtg tctaagaaca aaagcgaccc 4320
attcagagac tgcctctgaa acctagtact gcccacatg aggaaggaa agcaatggtg 4380
tcagtatcca ggccttctac agagtgttt tctgttagt tcttactttt tctgtttgt 4440
ttttttaaag atgaaataaa gaaccagggg gag 4470

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0210- 6

0211- 1255

0212- PRT

0213- Homo sapiens

0400- 6

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Met Glu Leu Ala Ala Leu Cys Arg Trp Gly Leu Leu Leu Ala Leu Leu
1          5          10          15
Pro Pro Gly Ala Ala Ser Thr Gln Val Cys Thr Gly Thr Asp Met Lys
20          25          30
Leu Arg Leu Pro Ala Ser Pro Glu Thr His Leu Asp Met Leu Arg His
35          40          45
Leu Tyr Gln Gly Cys Gln Val Val Gln Gly Asn Leu Glu Leu Thr Tyr
50          55          60
Leu Pro Thr Asn Ala Ser Leu Ser Phe Leu Gln Asp Ile Gln Glu Val
65          70          75          80

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Gln Gly Tyr Val Leu Ile Ala His Asn Gln Val Arg Gln Val Pro Leu  
 85 90 95  
 Gln Arg Leu Arg Ile Val Arg Gly Thr Gln Leu Phe Glu Asp Asn Tyr  
 100 105 110  
 Ala Leu Ala Val Leu Asp Asn Gly Asp Pro Leu Asn Asn Thr Thr Pro  
 115 120 125  
 Val Thr Gly Ala Ser Pro Gly Gly Leu Arg Glu Leu Gln Leu Arg Ser  
 130 135 140  
 Leu Thr Glu Ile Leu Lys Gly Gly Val Leu Ile Gln Arg Asn Pro Gln  
 145 150 155 160  
 Leu Cys Tyr Gln Asp Thr Ile Leu Trp Lys Asp Ile Phe His Lys Asn  
 165 170 175  
 Asn Gln Leu Ala Leu Thr Leu Ile Asp Thr Asn Arg Ser Arg Ala Cys  
 180 185 190  
 His Pro Cys Ser Pro Met Cys Lys Gly Ser Arg Cys Trp Gly Glu Ser  
 195 200 205  
 Ser Glu Asp Cys Gln Ser Leu Thr Arg Thr Val Cys Ala Gly Gly Cys  
 210 215 220  
 Ala Arg Cys Lys Gly Pro Leu Pro Thr Asp Cys Cys His Glu Gln Cys  
 225 230 235 240  
 Ala Ala Gly Cys Thr Gly Pro Lys His Ser Asp Cys Leu Ala Cys Leu  
 245 250 255  
 His Phe Asn His Ser Gly Ile Cys Glu Leu His Cys Pro Ala Leu Val  
 260 265 270  
 Thr Tyr Asn Thr Asp Thr Phe Glu Ser Met Pro Asn Pro Glu Gly Arg  
 275 280 285  
 Tyr Thr Phe Gly Ala Ser Cys Val Thr Ala Cys Pro Tyr Asn Tyr Leu  
 290 295 300  
 Ser Thr Asp Val Gly Ser Cys Thr Leu Val Cys Pro Leu His Asn Gln  
 305 310 315 320  
 Glu Val Thr Ala Glu Asp Gly Thr Gln Arg Cys Glu Lys Cys Ser Lys  
 325 330 335  
 Pro Cys Ala Arg Val Cys Tyr Gly Leu Gly Met Glu His Leu Arg Gln  
 340 345 350  
 Val Arg Ala Val Thr Ser Ala Asn Ile Gln Glu Phe Ala Gly Cys Lys  
 355 360 365  
 Lys Ile Phe Gly Ser Leu Ala Phe Leu Pro Glu Ser Phe Asp Gly Asp  
 370 375 380  
 Pro Ala Ser Asn Thr Ala Pro Leu Gln Pro Glu Gln Leu Gln Val Phe  
 385 390 395 400  
 Glu Thr Leu Glu Glu Ile Thr Gly Tyr Leu Tyr Ile Ser Ala Trp Pro  
 405 410 415  
 Asp Ser Leu Pro Asp Leu Ser Val Phe Gln Asn Leu Gln Val Ile Arg  
 420 425 430  
 Gly Arg Ile Leu His Asn Gly Ala Tyr Ser Leu Thr Leu Gln Gly Leu  
 435 440 445  
 Gly Ile Ser Trp Leu Gly Leu Arg Ser Leu Arg Gln Leu Gly Ser Gly  
 450 455 460  
 Leu Ala Leu Ile His His Asn Thr His Leu Cys Phe Val His Thr Val  
 465 470 475 480  
 Pro Trp Asp Gln Leu Phe Arg Asn Pro His Gln Ala Leu Leu His Thr  
 485 490 495  
 Ala Asn Arg Pro Glu Asp Glu Cys Val Gly Glu Gly Leu Ala Cys His  
 500 505 510

Gln Leu Cys Ala Arg Gly His Cys Trp Gly Pro Gly Pro Thr Gln Cys  
 515 520 525  
 Val Asn Cys Ser Gln Phe Leu Arg Gly Gln Glu Cys Val Glu Glu Cys  
 530 535 540  
 Arg Val Leu Gln Gly Leu Pro Arg Glu Tyr Val Asn Ala Arg His Cys  
 545 550 555 560  
 Leu Pro Cys His Pro Gly Cys Gln Pro Gln Asn Gly Ser Val Thr Cys  
 565 570 575  
 Phe Gly Pro Glu Ala Asp Gln Lys Val Ala Cys Ala His Tyr Lys Asp  
 580 585 590  
 Pro Pro Phe Cys Val Ala Arg Lys Pro Ser Gly Val Lys Pro Asp Leu  
 595 600 605  
 Ser Tyr Met Pro Ile Trp Lys Phe Pro Asp Glu Gln Gly Ala Cys Gln  
 610 615 620  
 Pro Lys Pro Ile Asn Cys Thr His Ser Cys Val Asp Leu Asp Asp Lys  
 625 630 635 640  
 Gly Cys Pro Ala Glu Gln Arg Ala Ser Pro Leu Thr Ser Ile Ile Ser  
 645 650 655  
 Ala Val Val Gly Ile Leu Leu Val Val Val Leu Gly Val Val Phe Gly  
 660 665 670  
 Ile Leu Ile Lys Arg Arg Gln Gln Lys Ile Arg Lys Tyr Thr Met Arg  
 675 680 685  
 Arg Leu Leu Gln Gly Thr Glu Leu Val Gly Pro Leu Thr Pro Ser Gly  
 690 695 700  
 Ala Met Pro Asn Gln Ala Gln Met Arg Ile Leu Lys Glu Thr Glu Leu  
 705 710 715 720  
 Arg Lys Val Lys Val Leu Gly Ser Gly Ala Phe Gly Thr Val Tyr Lys  
 725 730 735  
 Gly Ile Trp Ile Pro Asp Gly Gln Asn Val Lys Ile Pro Val Ala Leu  
 740 745 750  
 Lys Val Leu Arg Glu Asn Thr Ser Pro Lys Ala Asn Lys Glu Ile Leu  
 755 760 765  
 Asp Glu Ala Tyr Val Met Ala Gly Val Gly Ser Pro Tyr Val Ser Arg  
 770 775 780  
 Leu Leu Gly Ile Cys Leu Thr Ser Thr Val Gln Leu Val Thr Gln Leu  
 785 790 795 800  
 Met Pro Tyr Gly Cys Leu Leu Asp His Val Arg Gln Asn Arg Gly Arg  
 805 810 815  
 Leu Gly Ser Gln Asp Leu Leu Asn Trp Cys Met Gln Ile Ala Lys Gly  
 820 825 830  
 Met Ser Tyr Leu Glu Asp Val Arg Leu Val His Arg Asp Leu Ala Ala  
 835 840 845  
 Arg Asn Val Leu Val Lys Ser Pro Asn His Val Lys Ile Thr Asp Phe  
 850 855 860  
 Gly Leu Ala Arg Leu Leu Asp Ile Asp Glu Thr Glu Tyr His Ala Asp  
 865 870 875 880  
 Gly Gly Lys Val Pro Ile Lys Trp Met Ala Leu Gln Ser Ile Leu Arg  
 885 890 895  
 Arg Arg Phe Thr His Gln Ser Asp Val Trp Ser Tyr Gly Val Thr Val  
 900 905 910  
 Trp Gln Leu Met Thr Phe Gly Ala Lys Pro Tyr Asp Gly Ile Pro Ala  
 915 920 925  
 Arg Gln Ile Pro Asp Leu Leu Glu Lys Gly Glu Arg Leu Pro Gln Pro  
 930 935 940

Pro Ile Cys Thr Ile Asp Val Tyr Met Ile Met Val Lys Cys Trp Met  
 945 950 955 960  
 Ile Asp Ser Glu Cys Arg Pro Arg Phe Arg Glu Leu Val Ser Glu Phe  
 965 970 975  
 Ser Arg Met Ala Arg Asp Pro Gln Arg Phe Val Val Ile Gln Asn Glu  
 980 985 990  
 Asp Leu Gly Pro Ala Ser Pro Leu Asp Ser Thr Phe Tyr Arg Ser Leu  
 995 1000 1005  
 Leu Glu Asp Asp Asp Met Gly Asp Leu Val Asp Ala Glu Glu Tyr Leu  
 1010 1015 1020  
 Val Pro Gln Gln Gly Phe Phe Cys Pro Asp Pro Ala Pro Gly Ala Gly  
 1025 1030 1035 1040  
 Gly Met Val His His Arg His Arg Ser Ser Ser Thr Arg Ser Gly Gly  
 1045 1050 1055  
 Gly Asp Leu Thr Leu Gly Leu Gln Pro Ser Glu Gln Glu Ala Pro Arg  
 1060 1065 1070  
 Ser Pro Leu Ala Pro Ser Glu Gly Ala Gly Ser Asp Val Phe Asp Gly  
 1075 1080 1085  
 Asp Leu Gly Met Gly Ala Ala Lys Gly Leu Gln Ser Leu Pro Thr His  
 1090 1095 1100  
 Asp Pro Ser Pro Leu Gln Arg Tyr Ser Glu Asp Pro Thr Val Pro Leu  
 1105 1110 1115 1120  
 Pro Ser Glu Thr Asp Gly Tyr Val Ala Pro Leu Thr Cys Ser Pro Gln  
 1125 1130 1135  
 Pro Glu Tyr Val Asn Gln Pro Asp Val Arg Pro Gln Pro Pro Ser Pro  
 1140 1145 1150  
 Arg Glu Gly Pro Leu Pro Ala Ala Arg Pro Ala Gly Ala Thr Leu Glu  
 1155 1160 1165  
 Arg Pro Lys Thr Leu Ser Pro Gly Lys Asn Gly Val Val Lys Asp Val  
 1170 1175 1180  
 Phe Ala Phe Gly Gly Ala Val Glu Asn Pro Glu Tyr Leu Thr Pro Gln  
 1185 1190 1195 1200  
 Gly Gly Ala Ala Pro Gln Pro His Pro Pro Pro Ala Phe Ser Pro Ala  
 1205 1210 1215  
 Phe Asp Asn Leu Tyr Tyr Trp Asp Gln Asp Pro Pro Glu Arg Gly Ala  
 1220 1225 1230  
 Pro Pro Ser Thr Phe Lys Gly Thr Pro Thr Ala Glu Asn Pro Glu Tyr  
 1235 1240 1245  
 Leu Gly Leu Asp Val Pro Val  
 1250 1255

0210: 7

0211: 329

0212: INA

0213: *Trichophila melanogaster*

0400: 7

gcatttcag agggctacct caagtcctgc tcttcgact atctttcgga caactttgac	60
accgggtgt ttgtgggcac catcttcttt ttcagcttg tgtgtccac gctgatgac	120
ctttactact actgcagat cgtgggcac gtcttcagcc acgaaaagg cctacgggag	180
caggccaaga aaatgaacgt ggagtcgctg cctccaatg tggacaagag caaggagag	240
ggcagatag ggattgcgaa ggcgctatc accatctgt tctgttctt cgtgtcgtg	300
acgcctaag gcgtaatgtc gctgatcgg gattcgggg ataagagtct gcttacaaa	360
ggagccacga tgatcccggc ctgcacctgc aaactggtg cgtgcataga cccattcgtc	420

tatgcacataa gtcacccacag ataccggttg gagctgcaga agcgtgtgccc ctggctggga 480  
 gtcaacgaaa agtctgggga gatctcttcg gcgcagttca cgaaccacca ggagcagcaa 540  
 cagactaccg ctgcatagaa ccaaggacaa ctctactcta agacaactga ccattgtaaca 600  
 tgaagaccaa ggaaaaagta taaatgtccg acaacgaaac tgtataacat taattttata 660  
 attttagtgg tgacattctt gattttgaaa taaataaata gtaatttatt gcaaacgaag 720  
 tagaaaatg 729

0210 - 2

0211 - 373

0212 - PRT

0213 - *Drosophila melanogaster*

0400 - 3

Met Glu Pro Leu Cys Asn Ala Ser Glu Pro Pro Leu Arg Pro Glu Ala  
 1 5 10 15  
 Arg Ser Ser Gly Asn Gly Asp Leu Gln Phe Leu Gly Trp Asn Val Pro  
 20 25 30  
 Pro Asp Gln Ile Gln Tyr Ile Pro Glu His Trp Leu Thr Gln Leu Glu  
 35 40 45  
 Pro Pro Ala Ser Met His Tyr Met Leu Gly Val Phe Tyr Ile Phe Leu  
 50 55 60  
 Phe Cys Ala Ser Thr Val Gly Asn Gly Met Val Ile Trp Ile Phe Ser  
 65 70 75 80  
 Thr Ser Lys Ser Leu Arg Thr Pro Ser Asn Met Phe Val Leu Asn Leu  
 85 90 95  
 Ala Val Phe Asp Leu Ile Met Cys Leu Lys Ala Pro Ile Phe Asn Ser  
 100 105 110  
 Phe His Arg Gly Phe Ala Ile Tyr Leu Gly Asn Thr Trp Cys Gln Ile  
 115 120 125  
 Phe Ala Ser Ile Gly Ser Tyr Ser Gly Ile Gly Ala Gly Met Thr Asn  
 130 135 140  
 Ala Ala Ile Gly Tyr Asp Arg Tyr Asn Val Ile Thr Lys Pro Met Asn  
 145 150 155 160  
 Arg Asn Met Thr Phe Thr Lys Ala Val Ile Met Asn Ile Ile Ile Trp  
 165 170 175  
 Leu Tyr Cys Thr Pro Trp Val Val Leu Pro Leu Thr Gln Phe Trp Asp  
 180 185 190  
 Arg Phe Val Pro Glu Gly Tyr Leu Thr Ser Cys Ser Phe Asp Tyr Leu  
 195 200 205  
 Ser Asp Asn Phe Asp Thr Arg Leu Phe Val Gly Thr Ile Phe Phe Phe  
 210 215 220  
 Ser Phe Val Cys Pro Thr Leu Met Ile Leu Tyr Tyr Tyr Ser Gln Ile  
 225 230 235 240  
 Val Gly His Val Phe Ser His Glu Lys Ala Leu Arg Glu Gln Ala Lys  
 245 250 255  
 Lys Met Asn Val Glu Ser Leu Arg Ser Asn Val Asp Lys Ser Lys Glu  
 260 265 270  
 Thr Ala Glu Ile Arg Ile Ala Lys Ala Ala Ile Thr Ile Cys Phe Leu  
 275 280 285  
 Phe Phe Val Ser Trp Thr Pro Tyr Gly Val Met Ser Leu Ile Gly Ala  
 290 295 300  
 Phe Gly Asp Lys Ser Leu Leu Thr Gln Gly Ala Thr Met Ile Pro Ala  
 305 310 315 320  
 Cys Thr Cys Lys Leu Val Ala Cys Ile Asp Pro Phe Val Tyr Ala Ile

325                      330                      335  
 Ser His Pro Arg Tyr Arg Leu Glu Leu Gln Lys Arg Cys Pro Trp Leu  
                          340                      345                      350  
 Gly Val Asn Glu Lys Ser Gly Glu Ile Ser Ser Ala Gln Ser Thr Thr  
                          355                      360                      365  
 Thr Gln Glu Gln Gln Gln Thr Thr Ala Ala  
                          370                      375

#10 - 9  
 #11 - 191  
 #12 - PRT  
 #13 - Homo sapiens

#400 - 9  
 Met Glu Leu Ala Ala Leu Cys Arg Trp Gly Leu Leu Leu Ala Leu Leu  
   1                      5                      10                      15  
 Pro Pro Gly Ala Ala Ser Thr Gln Val Cys Thr Gly Thr Asp Met Lys  
                          20                      25                      30  
 Leu Arg Leu Pro Ala Ser Pro Glu Thr His Leu Asp Met Leu Arg His  
                          35                      40                      45  
 Leu Tyr Gln Gly Cys Gln Val Val Gln Gly Asn Leu Glu Leu Thr Tyr  
   50                      55                      60  
 Leu Pro Thr Asn Ala Ser Leu Ser Phe Leu Gln Asp Ile Gln Glu Val  
   65                      70                      75                      80  
 Gln Gly Tyr Val Leu Ile Ala His Asn Gln Val Arg Gln Val Pro Leu  
                          85                      90                      95  
 Gln Arg Leu Arg Ile Val Arg Gly Thr Gln Leu Phe Glu Asp Asn Tyr  
                          100                      105                      110  
 Ala Leu Ala Val Leu Asp Asn Gly Asp Pro Leu Asn Asn Thr Thr Pro  
                          115                      120                      125  
 Val Thr Gly Ala Ser Pro Gly Gly Leu Arg Glu Leu Gln Leu Arg Ser  
                          130                      135                      140  
 Leu Thr Glu Ile Leu Lys Gly Gly Val Leu Ile Gln Arg Asn Pro Gln  
   145                      150                      155                      160  
 Leu Cys Tyr Gln Asp Thr Ile Leu Trp Lys Asp Ile Phe His Lys Asn  
                          165                      170                      175  
 Asn Gln Leu Ala Leu Thr Leu Ile Asp Thr Asn Arg Ser Arg Ala  
                          180                      185                      190

#10 - 10  
 #11 - 1277  
 #12 - DNA  
 #13 - Homo sapiens

#400 - 10  
 gtcatctgct atttttaaac ttccctggaa taatatatgt aatctacttc taataagttt    60  
 ttcttattca gcatttttgt ttaaactaat ttataattat ttageccttat ttctccatgt    120  
 ttaacttgct ttaaagctca gcactgggtgt ttccagccat ggctctccca ttttaaggct    180  
 attttaattc atttattatt ctggaatata tcccttaata atttatttag gaaggctgtc    240  
 tggtaggtgg tatttctgtt gcagttgttg tttctctgcc tggtaggtga catatttcta    300  
 ttgacttgac acttaactgg catcttatct aggtagataa tgctaattca aaattctgca    360  
 gatattggtc tgttgttttt tgccatttat ggtagagtaa gatgccaaagt tggtttttgg    420  
 ttctctgtag tcattctgtt ttcatattgt ttttagcttt gcctttggaa tttaaaatgt    480  
 tcaaaatgat ttgtctggat gagaatogat tttcataact tttgctttga tacactaaac    540



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agtttgagtt tctagatgat gccattttta attcatacga ggaaatatct tctagtatag      600
ttttctgcttg attaattcta tgtttgtctc tttagggacat ctattaatttt tataatgctg      650
ccttttttttc agactttctgt ttcagaatat togttttctat gaatgtaate cttggctata      700
gtaggaatga aataataaaa gcagtagctt ctgtctgccc tcccttggtta tgcagtcctt      750
acagacatto tccccaccto ccatccccc accccagctc agtgaaactc tccacacttt      800
ggttctggaa attggcaggy ttaggttggc actcactccc aatccacato cacaataaat      850
cactttttat tatcttatca aaatctgtag aatgcctctt tattctattt tgttctgtcg      900
gaggttttgtt tttcttttct aattatttta tttcttaggt tttttgaggy aatttcaaga      950
ggggagatttt tttattcagg ctcatcttaa cgtcatgtct ggaactcaag ctactgaatt      1000
atatattctt taatacatat agacttaagt caatgagttt aaactgcaag gaaagtggtta      1050
aatttcttccc tcaagtgttg tcaaaatctg tagagaaaag aggaacagct tctcttaag      1100
aaagttagct ggttaggtat acagtcattg ccgaggaagg cttgcacagg gtgaaagctt      1150
tgcctctctg ctgtctg      1177

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OR10: 11

OR11: 356

OR12: PRT

OR13: Mus musculus

OR20: 11

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Met Ala Lys Thr Ile Arg Arg Leu Ser Val Ala Phe Leu Thr Leu Ser
1      5      10      15
Asp Arg Gly Pro His Tyr Lys Ile Leu Leu Pro Leu Pro His Lys Gly
20     25     30
Trp Thr Pro Gly Leu Thr His Asn Ala Ser Leu Tyr Cys Ala Ser Ile
35     40     45
Ile Leu Lys Asn Thr Met Gly Leu Ala Ile Leu Ile Phe Val Thr Val
50     55     60
Leu Leu Ile Ser Asp Ala Val Ser Val Glu Thr Gln Ala Tyr Phe Asn
65     70     75     80
Gly Thr Ala Tyr Leu Pro Cys Pro Phe Thr Lys Ala Gln Asn Ile Ser
85     90     95
Leu Ser Glu Leu Val Val Phe Trp Gln Asp Gln Gln Lys Leu Val Leu
100    105    110
Tyr Glu His Tyr Leu Gly Thr Glu Lys Leu Asp Ser Val Asn Ala Lys
115    120    125
Tyr Leu Gly Arg Thr Ser Phe Asp Arg Asn Asn Trp Thr Leu Arg Leu
130    135    140
His Asn Val Gln Ile Lys Asp Met Gly Ser Tyr Asp Cys Phe Ile Gln
145    150    155    160
Lys Lys Pro Pro Thr Gly Ser Ile Ile Leu Gln Gln Thr Leu Thr Glu
165    170    175
Leu Ser Val Ile Ala Asn Phe Ser Glu Pro Glu Ile Lys Leu Ala Gln
180    185    190
Asn Val Thr Gly Asn Ser Gly Ile Asn Leu Thr Cys Thr Ser Lys Gln
195    200    205
Gly His Pro Lys Pro Lys Lys Met Tyr Phe Leu Ile Thr Asn Ser Thr
210    215    220
Asn Glu Tyr Gly Asp Asn Met Gln Ile Ser Gln Asp Asn Val Thr Glu
225    230    235    240
Leu Phe Ser Ile Ser Asn Ser Leu Ser Leu Ser Phe Pro Asp Gly Val
245    250    255
Trp His Met Thr Val Val Cys Val Leu Glu Thr Glu Ser Met Lys Ile
260    265    270

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Ser Ser Lys Pro Leu Asn Phe Thr Gln Glu Phe Pro Ser Pro Gln Thr  
 275 280 285  
 Tyr Trp Lys Glu Ile Thr Ala Ser Val Thr Val Ala Leu Leu Leu Val  
 290 295 300  
 Met Leu Leu Ile Ile Val Lys His Lys Lys Pro Asn Gln Pro Ser Arg  
 305 310 315 320  
 Pro Ser Asn Thr Ala Ser Lys Leu Glu Arg Asp Ser Asn Ala Asp Arg  
 325 330 335  
 Glu Thr Ile Asn Leu Lys Glu Leu Glu Pro Gln Ile Ala Ser Ala Lys  
 340 345 350  
 Pro Asn Ala Glu  
 355

0010: 12

0011: 356

0012: PRT

0013: Mus musculus

0400: 12

Met Ala Lys Thr Ile Arg Arg Leu Ser Val Ala Phe Leu Thr Leu Ser  
 1 5 10 15  
 Asp Arg Gly Pro His Tyr Lys Ile Leu Pro Leu Pro His Lys Gly  
 20 25 30  
 Pro Thr Pro Gly Leu Thr His Asn Ala Ser Leu Tyr Lys Ala Ser Ile  
 35 40 45  
 Ile Leu Lys Asn Thr Met Gly Leu Ala Ile Leu Ile Phe Val Thr Val  
 50 55 60  
 Leu Leu Ile Ser Asp Ala Val Ser Val Glu Thr Gln Ala Tyr Phe Asn  
 65 70 75 80  
 Gly Thr Ala Tyr Leu Pro Lys Pro Phe Thr Lys Ala Gln Asn Ile Ser  
 85 90 95  
 Leu Ser Glu Leu Val Val Phe Trp Gln Asp Gln Gln Lys Leu Val Leu  
 100 105 110  
 Tyr Glu His Tyr Leu Gly Thr Glu Lys Leu Asp Ser Val Asn Ala Lys  
 115 120 125  
 Tyr Leu Gly Arg Thr Ser Phe Asp Arg Asn Asn Trp Thr Leu Arg Leu  
 130 135 140  
 His Asn Val Gln Ile Lys Asp Met Gly Ser Tyr Asp Cys Phe Ile Gln  
 145 150 155 160  
 Lys Lys Pro Pro Thr Gly Ser Ile Ile Leu Gln Gln Thr Leu Thr Glu  
 165 170 175  
 Leu Ser Val Ile Ala Asn Phe Ser Glu Pro Glu Ile Lys Leu Ala Gln  
 180 185 190  
 Asn Val Thr Gly Asn Ser Gly Ile Asn Leu Thr Cys Thr Ser Lys Gln  
 195 200 205  
 Gly His Pro Lys Pro Lys Lys Met Tyr Phe Leu Ile Thr Asn Ser Thr  
 210 215 220  
 Asn Glu Tyr Gly Asp Asn Met Gln Ile Ser Gln Asp Asn Val Thr Glu  
 225 230 235 240  
 Leu Phe Ser Ile Ser Asn Ser Leu Ser Leu Ser Phe Pro Asp Gly Val  
 245 250 255  
 Trp His Met Thr Val Val Cys Val Leu Glu Thr Glu Ser Met Lys Ile  
 260 265 270  
 Ser Ser Lys Pro Leu Asn Phe Thr Gln Glu Phe Pro Ser Pro Gln Thr

275                      280                      285  
 Tyr Trp Lys Glu Ile Thr Ala Ser Val Thr Val Ala Leu Leu Leu Val  
 290                      295                      300  
 Met Leu Leu Ile Ile Val Cys His Lys Lys Pro Asn Gln Pro Ser Arg  
 305                      310                      315                      320  
 Pro Ser Asn Thr Ala Ser Lys Leu Glu Arg Asp Ser Asn Ala Asp Arg  
 325                      330                      335  
 Glu Thr Ile Asn Leu Lys Glu Leu Glu Pro Gln Ile Ala Ser Ala Lys  
 340                      345                      350  
 Pro Asn Ala Glu  
 355

CL10 - 13

CL11 - 309

CL12 - PRF

CL13 - Mus musculus

CG00 - 13

Met Asp Pro Arg Cys Thr Met Gly Leu Ala Ile Leu Ile Phe Val Thr  
 1                      5                      10                      15  
 Val Leu Leu Ile Ser Asp Ala Val Ser Val Glu Thr Gln Ala Tyr Phe  
 20                      25                      30  
 Asn Gly Thr Ala Tyr Leu Pro Cys Pro Phe Thr Lys Ala Gln Asn Ile  
 35                      40                      45  
 Ser Leu Ser Glu Leu Val Val Phe Trp Gln Asp Gln Gln Lys Leu Val  
 50                      55                      60  
 Leu Tyr Glu His Tyr Leu Gly Thr Glu Lys Leu Asp Ser Val Asn Ala  
 65                      70                      75                      80  
 Lys Tyr Leu Gly Arg Thr Ser Phe Asp Arg Asn Asn Trp Thr Leu Arg  
 85                      90                      95  
 Leu His Asn Val Gln Ile Lys Asp Met Gly Ser Tyr Asp Cys Phe Ile  
 100                      105                      110  
 Gln Lys Lys Pro Pro Thr Gly Ser Ile Ile Leu Gln Gln Thr Leu Thr  
 115                      120                      125  
 Glu Leu Ser Val Ile Ala Asn Phe Ser Glu Pro Glu Ile Lys Leu Ala  
 130                      135                      140  
 Gln Asn Val Thr Gly Asn Ser Gly Ile Asn Leu Thr Cys Thr Ser Lys  
 145                      150                      155                      160  
 Gln Gly His Pro Lys Pro Lys Lys Met Tyr Phe Leu Ile Thr Asn Ser  
 165                      170                      175  
 Thr Asn Glu Tyr Gly Asp Asn Met Gln Ile Ser Gln Asp Asn Val Thr  
 180                      185                      190  
 Glu Leu Phe Ser Ile Ser Asn Ser Leu Ser Leu Ser Phe Pro Asp Gly  
 195                      200                      205  
 Val Trp His Met Thr Val Val Cys Val Leu Glu Thr Glu Ser Met Lys  
 210                      215                      220  
 Ile Ser Ser Lys Pro Leu Asn Phe Thr Gln Glu Phe Pro Ser Pro Gln  
 225                      230                      235                      240  
 Thr Tyr Trp Lys Glu Ile Thr Ala Ser Val Thr Val Ala Leu Leu Leu  
 245                      250                      255  
 Val Met Leu Leu Ile Ile Val Cys His Lys Lys Pro Asn Gln Pro Ser  
 260                      265                      270  
 Arg Pro Ser Asn Thr Ala Ser Lys Leu Glu Arg Asp Ser Asn Ala Asp  
 275                      280                      285

Arg Glu Thr Ile Asn Leu Lys Glu Leu Glu Pro Gln Ile Ala Ser Ala  
 230 295 300

Lys Pro Asn Ala Glu  
 305

0210: 14

0211: 314

0212: PRT

0213: Mus musculus

0400: 14

Met Tyr Val Ile Lys Thr Cys Ala Thr Cys Thr Met Gly Leu Ala Ile  
 1 5 10 15  
 Leu Ile Phe Val Thr Val Leu Leu Ile Ser Asp Ala Val Ser Val Glu  
 20 25 30  
 Thr Gln Ala Tyr Phe Asn Gly Thr Ala Tyr Leu Pro Lys Pro Phe Thr  
 35 40 45  
 Lys Ala Gln Asn Ile Ser Leu Ser Glu Leu Val Val Phe Trp Gln Asp  
 50 55 60  
 Gln Gln Lys Leu Val Leu Tyr Glu His Tyr Leu Gly Thr Glu Lys Leu  
 65 70 75 80  
 Asp Ser Val Asn Ala Lys Tyr Leu Gly Arg Thr Ser Phe Asp Arg Asn  
 85 90 95  
 Asn Trp Thr Leu Arg Leu His Asn Val Gln Ile Lys Asp Met Gly Ser  
 100 105 110  
 Tyr Asp Cys Phe Ile Gln Lys Lys Pro Pro Thr Gly Ser Ile Ile Leu  
 115 120 125  
 Gln Gln Thr Leu Thr Glu Leu Ser Val Ile Ala Asn Phe Ser Glu Pro  
 130 135 140  
 Glu Ile Lys Leu Ala Gln Asn Val Thr Gly Asn Ser Gly Ile Asn Leu  
 145 150 155 160  
 Thr Lys Thr Ser Lys Gln Gly His Pro Lys Pro Lys Lys Met Tyr Phe  
 165 170 175  
 Leu Ile Thr Asn Ser Thr Asn Glu Tyr Gly Asp Asn Met Gln Ile Ser  
 180 185 190  
 Gln Asp Asn Val Thr Glu Leu Phe Ser Ile Ser Asn Ser Leu Ser Leu  
 195 200 205  
 Ser Ile Pro Asp Gly Val Trp His Met Thr Val Val Cys Val Leu Glu  
 210 215 220  
 Thr Glu Ser Met Lys Ile Ser Ser Lys Pro Leu Asn Phe Thr Gln Glu  
 225 230 235 240  
 Phe Pro Ser Pro Gln Thr Tyr Trp Lys Glu Ile Thr Ala Ser Val Thr  
 245 250 255  
 Val Ala Leu Leu Leu Val Met Leu Leu Ile Ile Val Cys His Lys Lys  
 260 265 270  
 Pro Asn Gln Pro Ser Arg Pro Ser Asn Thr Ala Ser Lys Leu Glu Arg  
 275 280 285  
 Asp Ser Asn Ala Asp Arg Glu Thr Ile Asn Leu Lys Glu Leu Glu Pro  
 290 295 300  
 Gln Ile Ala Ser Ala Lys Pro Asn Ala Glu  
 305 310

0210: 15

0211: 303

0212: PRT

0213: Mus musculus

0400: 15

Met	Gly	Leu	Ala	Ile	Leu	Ile	Phe	Val	Thr	Val	Leu	Leu	Ile	Ser	Asp
1				5					10					15	
Ala	Val	Ser	Val	Glu	Thr	Gln	Ala	Tyr	Phe	Asn	Gly	Thr	Ala	Tyr	Leu
			20					25					30		
Pro	Lys	Pro	Phe	Thr	Lys	Ala	Gln	Asn	Ile	Ser	Leu	Ser	Glu	Leu	Val
		35					40					45			
Val	Phe	Trp	Gln	Asp	Gln	Gln	Lys	Leu	Val	Leu	Tyr	Glu	His	Tyr	Leu
	50					55					60				
Gly	Thr	Glu	Lys	Leu	Asp	Ser	Val	Asn	Ala	Lys	Tyr	Leu	Gly	Arg	Thr
65					70					75				80	
Ser	Phe	Asp	Arg	Asn	Asn	Trp	Thr	Leu	Arg	Leu	His	Asn	Val	Gln	Ile
				85					90					95	
Lys	Asp	Met	Gly	Ser	Tyr	Asp	Cys	Phe	Ile	Gln	Lys	Lys	Pro	Pro	Thr
		100						105					110		
Gly	Ser	Ile	Ile	Leu	Gln	Gln	Thr	Leu	Thr	Glu	Leu	Ser	Val	Ile	Ala
	115						120					125			
Asn	Phe	Ser	Glu	Pro	Glu	Ile	Lys	Leu	Ala	Gln	Asn	Val	Thr	Gly	Asn
	130					135					140				
Ser	Gly	Ile	Asn	Leu	Thr	Cys	Thr	Ser	Lys	Gln	Gly	His	Pro	Lys	Pro
145					150					155				160	
Lys	Lys	Met	Tyr	Phe	Leu	Ile	Thr	Asn	Ser	Thr	Asn	Glu	Tyr	Gly	Asp
			165						170					175	
Asn	Met	Gln	Ile	Ser	Gln	Asp	Asn	Val	Thr	Glu	Leu	Phe	Ser	Ile	Ser
	180							185					190		
Asn	Ser	Leu	Ser	Leu	Ser	Phe	Pro	Asp	Gly	Val	Trp	His	Met	Thr	Val
	195						200					205			
Val	Cys	Val	Leu	Glu	Thr	Glu	Ser	Met	Lys	Ile	Ser	Ser	Lys	Pro	Leu
	210					215					220				
Asn	Phe	Thr	Gln	Glu	Phe	Pro	Ser	Pro	Gln	Thr	Tyr	Trp	Lys	Glu	Ile
225				230						235				240	
Thr	Ala	Ser	Val	Thr	Val	Ala	Leu	Leu	Leu	Val	Met	Leu	Leu	Ile	Ile
			245							250				255	
Val	Cys	His	Lys	Lys	Pro	Asn	Gln	Pro	Ser	Arg	Pro	Ser	Asn	Thr	Ala
			260					265				270			
Ser	Lys	Leu	Glu	Arg	Asp	Ser	Asn	Ala	Asp	Arg	Glu	Thr	Ile	Asn	Leu
	275						280					285			
Lys	Glu	Leu	Glu	Pro	Gln	Ile	Ala	Ser	Ala	Lys	Pro	Asn	Ala	Glu	
	290					295						300			

0210: 16

0211: 356

0212: PRT

0213: Mus musculus

0400: 16

Met	Ala	Lys	Thr	Ile	Arg	Arg	Leu	Ser	Val	Ala	Phe	Leu	Thr	Leu	Ser
1				5					10					15	
Asp	Arg	Gly	Pro	His	Tyr	Lys	Ile	Leu	Leu	Pro	Leu	Pro	His	Lys	Gly
		20						25					30		
Trp	Thr	Pro	Gly	Leu	Thr	His	Asn	Ala	Ser	Leu	Tyr	Cys	Ala	Ser	Ile

35 40 45  
 Ile Leu Lys Asn Thr Met Gly Leu Ala Ile Leu Ile Phe Val Thr Val  
 50 55 60  
 Leu Leu Ile Ser Asp Ala Val Ser Val Glu Thr Gln Ala Tyr Phe Asn  
 65 70 75 80  
 Gly Thr Ala Tyr Leu Pro Cys Pro Phe Thr Lys Ala Gln Asn Ile Ser  
 85 90 95  
 Leu Ser Gln Leu Val Val Phe Trp Gln Asp Gln Gln Lys Leu Val Leu  
 100 105 110  
 Tyr Glu His Tyr Leu Gly Thr Glu Lys Leu Asp Ser Val Asn Ala Lys  
 115 120 125  
 Tyr Leu Gly Arg Thr Ser Phe Asp Arg Asn Asn Trp Thr Leu Arg Leu  
 130 135 140  
 His Asn Val Gln Ile Lys Asp Met Gly Ser Tyr Asp Cys Phe Ile Gln  
 145 150 155 160  
 Lys Lys Pro Pro Thr Gly Ser Ile Ile Leu Gln Gln Thr Leu Thr Gln  
 165 170 175  
 Leu Ser Val Ile Ala Asn Phe Ser Glu Pro Gln Ile Lys Leu Ala Gln  
 180 185 190  
 Asn Val Thr Gly Asn Ser Gly Ile Asn Leu Thr Cys Thr Ser Lys Gln  
 195 200 205  
 Gly His Pro Lys Pro Lys Lys Met Tyr Phe Leu Ile Thr Asn Ser Thr  
 210 215 220  
 Asn Glu Tyr Gly Asp Asn Met Gln Ile Ser Gln Asp Asn Val Thr Glu  
 225 230 235 240  
 Leu Phe Ser Ile Ser Asn Ser Leu Ser Leu Ser Phe Pro Asp Gly Val  
 245 250 255  
 Trp His Met Thr Val Val Cys Val Leu Glu Thr Glu Ser Met Lys Ile  
 260 265 270  
 Ser Ser Lys Pro Leu Asn Phe Thr Gln Glu Phe Pro Ser Pro Gln Thr  
 275 280 285  
 Tyr Trp Lys Gln Ile Thr Ala Ser Val Thr Val Ala Leu Leu Leu Val  
 290 295 300  
 Met Leu Leu Ile Ile Val Cys His Lys Lys Pro Asn Gln Pro Ser Arg  
 305 310 315 320  
 Pro Ser Asn Thr Ala Ser Lys Leu Glu Arg Asp Ser Asn Ala Asp Arg  
 325 330 335  
 Glu Thr Ile Asn Leu Lys Glu Leu Glu Pro Gln Ile Ala Ser Ala Lys  
 340 345 350  
 Pro Asn Ala Glu  
 355

4110: 17

4111: 355

4112: PRT

4113: Mus musculus

4400: 17

Met Ala Lys Thr Ile Arg Arg Leu Ser Val Ala Phe Leu Thr Leu Ser  
 1 5 10 15  
 Asp Arg Gly Pro His Tyr Lys Ile Leu Leu Pro Leu Pro His Lys Gly  
 20 25 30  
 Trp Thr Pro Gly Leu Thr His Asn Ala Ser Leu Tyr Cys Ala Ser Ile  
 35 40 45

Ile Leu Lys Asn Thr Met Gly Leu Ala Ile Leu Ile Phe Val Thr Val  
 50 55 60  
 Leu Leu Ile Ser Asp Ala Val Ser Val Glu Thr Gln Ala Tyr Phe Asn  
 65 70 75 80  
 Gly Thr Ala Tyr Leu Pro Cys Pro Phe Thr Lys Ala Gln Asn Ile Ser  
 85 90 95  
 Leu Ser Glu Leu Val Val Phe Trp Gln Asp Gln Gln Lys Leu Val Leu  
 100 105 110  
 Tyr Glu His Tyr Leu Gly Thr Glu Lys Leu Asp Ser Val Asn Ala Lys  
 115 120 125  
 Tyr Leu Gly Arg Thr Ser Phe Asp Arg Asn Asn Trp Thr Leu Arg Leu  
 130 135 140  
 His Asn Val Gln Ile Lys Asp Met Gly Ser Tyr Asp Cys Phe Ile Gln  
 145 150 155 160  
 Lys Lys Pro Pro Thr Gly Ser Ile Ile Leu Gln Gln Thr Leu Thr Glu  
 165 170 175  
 Leu Ser Val Ile Ala Asn Phe Ser Glu Pro Glu Ile Lys Leu Ala Gln  
 180 185 190  
 Asn Val Thr Gly Asn Ser Gly Ile Asn Leu Thr Cys Thr Ser Lys Gln  
 195 200 205  
 Gly His Pro Lys Pro Lys Lys Met Tyr Phe Leu Ile Thr Asn Ser Thr  
 210 215 220  
 Asn Glu Tyr Gly Asp Asn Met Gln Ile Ser Gln Asp Asn Val Thr Glu  
 225 230 235 240  
 Leu Phe Ser Ile Ser Asn Ser Leu Ser Leu Ser Phe Pro Asp Gly Val  
 245 250 255  
 Trp His Met Thr Val Val Cys Val Leu Glu Thr Glu Ser Met Lys Ile  
 260 265 270  
 Ser Ser Lys Pro Leu Asn Phe Thr Gln Glu Phe Pro Ser Pro Gln Thr  
 275 280 285  
 Tyr Trp Lys Glu Ile Thr Ala Ser Val Thr Val Ala Leu Leu Leu Val  
 290 295 300  
 Met Leu Leu Ile Ile Val Cys His Lys Lys Pro Asn Gln Pro Ser Arg  
 305 310 315 320  
 Pro Ser Asn Thr Ala Ser Lys Leu Glu Arg Asp Ser Asn Ala Asp Arg  
 325 330 335  
 Glu Thr Ile Asn Leu Lys Glu Leu Glu Pro Gln Ile Ala Ser Ala Lys  
 340 345 350  
 Pro Asn Ala Glu  
 355

-210- 18

-211- 309

-212- FRT

-213- Mus musculus

-400- 18

Met Asp Pro Arg Cys Thr Met Gly Leu Ala Ile Leu Ile Phe Val Thr  
 1 5 10 15  
 Val Leu Leu Ile Ser Asp Ala Val Ser Val Glu Thr Gln Ala Tyr Phe  
 20 25 30  
 Asn Gly Thr Ala Tyr Leu Pro Cys Pro Phe Thr Lys Ala Gln Asn Ile  
 35 40 45  
 Ser Leu Ser Glu Leu Val Val Phe Trp Gln Asp Gln Gln Lys Leu Val

50		55		60
Leu Tyr Glu His Tyr	Leu Gly Thr Glu Lys	Leu Asp Ser Val Asn Ala		
65	70	75	80	
Lys Tyr Leu Gly Arg Thr Ser Phe Asp Arg Asn Asn Trp Thr Leu Arg				
	85	90	95	
Leu His Asn Val Gln Ile Lys Asp Met Gly Ser Tyr Asp Cys Phe Ile				
	100	105	110	
Gln Lys Lys Pro Pro Thr Gly Ser Ile Ile Leu Gln Gln Thr Leu Thr				
	115	120	125	
Glu Leu Ser Val Ile Ala Asn Phe Ser Glu Pro Glu Ile Lys Leu Ala				
	130	135	140	
Gln Asn Val Thr Gly Asn Ser Gly Ile Asn Leu Thr Cys Thr Ser Lys				
145	150	155	160	
Gln Gly His Pro Lys Pro Lys Lys Met Tyr Phe Leu Ile Thr Asn Ser				
	165	170	175	
Thr Asn Glu Tyr Gly Asp Asn Met Gln Ile Ser Gln Asp Asn Val Thr				
	180	185	190	
Glu Leu Phe Ser Ile Ser Asn Ser Leu Ser Leu Ser Phe Pro Asp Gly				
	195	200	205	
Val Trp His Met Thr Val Val Cys Val Leu Glu Thr Glu Ser Met Lys				
	210	215	220	
Ile Ser Ser Lys Pro Leu Asn Phe Thr Gln Glu Phe Pro Ser Pro Gln				
225	230	235	240	
Thr Tyr Trp Lys Glu Ile Thr Ala Ser Val Thr Val Ala Leu Leu Leu				
	245	250	255	
Val Met Leu Leu Ile Ile Val Cys His Lys Lys Pro Asn Gln Pro Ser				
	260	265	270	
Arg Pro Ser Asn Thr Ala Ser Lys Leu Glu Arg Asp Ser Asn Ala Asp				
	275	280	285	
Arg Glu Thr Ile Asn Leu Lys Glu Leu Glu Pro Gln Ile Ala Ser Ala				
	290	295	300	
Lys Pro Asn Ala Glu				
305				

EF10 - 19

EF11 - 314

EF12 - PRT

EF13 - Mus musculus

EF400 - 19

Met Tyr Val Ile Lys Thr Cys Ala Thr Cys Thr Met Gly Leu Ala Ile				
1	5	10	15	
Leu Ile Phe Val Thr Val Leu Leu Ile Ser Asp Ala Val Ser Val Glu				
	20	25	30	
Thr Gln Ala Tyr Phe Asn Gly Thr Ala Tyr Leu Pro Cys Pro Phe Thr				
	35	40	45	
Lys Ala Gln Asn Ile Ser Leu Ser Glu Leu Val Val Phe Trp Gln Asp				
	50	55	60	
Gln Gln Lys Leu Val Leu Tyr Glu His Tyr Leu Gly Thr Glu Lys Leu				
65	70	75	80	
Asp Ser Val Asn Ala Lys Tyr Leu Gly Arg Thr Ser Phe Asp Arg Asn				
	85	90	95	
Asn Trp Thr Leu Arg Leu His Asn Val Gln Ile Lys Asp Met Gly Ser				
	100	105	110	



Tyr Asp Cys Phe Ile Gln Lys Lys Pro Pro Thr Gly Ser Ile Ile Leu  
 115 170 175  
 Gln Gln Thr Leu Thr Glu Leu Ser Val Ile Ala Asn Phe Ser Glu Pro  
 135 140  
 Glu Ile Lys Leu Ala Gln Asn Val Thr Gly Asn Ser Gly Ile Asn Leu  
 145 150 155 160  
 Thr Cys Thr Ser Lys Gln Gly His Pro Lys Pro Lys Lys Met Tyr Phe  
 165 170 175  
 Leu Ile Thr Asn Ser Thr Asn Glu Tyr Gly Asp Asn Met Gln Ile Ser  
 180 185 190  
 Gln Asp Asn Val Thr Glu Leu Phe Ser Ile Ser Asn Ser Leu Ser Leu  
 195 200 205  
 Ser Phe Pro Asp Gly Val Trp His Met Thr Val Val Lys Val Leu Glu  
 210 215 220  
 Thr Glu Ser Met Lys Ile Ser Ser Lys Pro Leu Asn Phe Thr Gln Glu  
 225 230 235 240  
 Phe Pro Ser Pro Gln Thr Tyr Trp Lys Glu Ile Thr Ala Ser Val Thr  
 245 250 255  
 Val Ala Leu Leu Leu Val Met Leu Leu Ile Ile Val Lys His Lys Lys  
 260 265 270  
 Pro Asn Gln Pro Ser Arg Pro Ser Asn Thr Ala Ser Lys Leu Glu Arg  
 275 280 285  
 Asp Ser Asn Ala Asp Arg Glu Thr Ile Asn Leu Lys Glu Leu Glu Pro  
 290 295 300  
 Gln Ile Ala Ser Ala Lys Pro Asn Ala Glu  
 305 310

0210 - 10

0211 - 203

0212 - PRF

0213 - Mus musculus

0400 - 10

Met Gly Leu Ala Ile Leu Ile Phe Val Thr Val Leu Leu Ile Ser Asp  
 1 5 10 15  
 Ala Val Ser Val Glu Thr Gln Ala Tyr Phe Asn Gly Thr Ala Tyr Leu  
 20 25 30  
 Pro Cys Pro Phe Thr Lys Ala Gln Asn Ile Ser Leu Ser Glu Leu Val  
 35 40 45  
 Val Phe Trp Gln Asp Gln Gln Lys Leu Val Leu Tyr Glu His Tyr Leu  
 50 55 60  
 Gly Thr Glu Lys Leu Asp Ser Val Asn Ala Lys Tyr Leu Gly Arg Thr  
 65 70 75 80  
 Ser Phe Asp Arg Asn Asn Trp Thr Leu Arg Leu His Asn Val Gln Ile  
 85 90 95  
 Lys Asp Met Gly Ser Tyr Asp Cys Phe Ile Gln Lys Lys Pro Pro Thr  
 100 105 110  
 Gly Ser Ile Ile Leu Gln Gln Thr Leu Thr Glu Leu Ser Val Ile Ala  
 115 120 125  
 Asn Phe Ser Glu Pro Glu Ile Lys Leu Ala Gln Asn Val Thr Gly Asn  
 130 135 140  
 Ser Gly Ile Asn Leu Thr Cys Thr Ser Lys Gln Gly His Pro Lys Pro  
 145 150 155 160  
 Lys Lys Met Tyr Phe Leu Ile Thr Asn Ser Thr Asn Glu Tyr Gly Asp

	165		170		175
Asn Met Gln Ile Ser Gln Asp Asn Val Thr Glu Leu Phe Ser Ile Ser					
	180		185		190
Asn Ser Leu Ser Leu Ser Phe Pro Asp Gly Val Trp His Met Thr Val					
	195		200		205
Val Cys Val Leu Glu Thr Glu Ser Met Lys Ile Ser Ser Lys Pro Leu					
	210		215		220
Asn Phe Thr Gln Glu Phe Pro Ser Pro Gln Thr Tyr Trp Lys Glu Ile					
	225		230		235
Thr Ala Ser Val Thr Val Ala Leu Leu Leu Val Met Leu Leu Ile Ile					
	245		250		255
Val Cys His Lys Lys Pro Asn Gln Pro Ser Arg Pro Ser Asn Thr Ala					
	260		265		270
Ser Lys Leu Glu Arg Asp Ser Asn Ala Asp Arg Glu Thr Ile Asn Leu					
	275		280		285
Lys Glu Leu Glu Pro Gln Ile Ala Ser Ala Lys Pro Asn Ala Glu					
	290		295		300

SI10: 21

SI11: 1424

SI12: DNA

SI13: Homo sapiens

SI400: 21

aggagcctta	ggaggtacgg	ggagctcgca	aatactcctt	ttggttttatt	cttaccacct	50
agcttcctgg	tcctctggga	atgctgctgt	gottatgpat	ctggctctctt	tttggagcta	100
cagtggaacg	gcatttggtg	cagcaactatg	ggactgagta	acattctctt	tgtgatggcc	150
ctcttgctct	ctgggtgctg	tcctctgaag	attcaagctt	atttcaatga	gactgcagac	200
ctgcaatgpc	aattttgaaa	ctctcaaaaac	caaagcctga	gtgagctagt	agtatttttg	250
caggacacgg	aaaacttggg	tctgaatgag	gtatacttag	gcacagagaa	atttgacagt	300
gtctcttcca	agtatatggg	ccgcacaagt	tttgattcgg	acagttggac	cctgagactt	350
caaatctctc	agatcaagga	caaggccttg	tatcaatgta	tcctccatca	caaaaagccc	400
acaggaatga	ttcgpatcca	ccagatgaat	tctgaactgt	cagtgccttg	taacttcagt	450
caacttgaaa	tagtaccaat	ttctaatata	acagaaaatg	tgtacataaa	tttgacctgc	500
ctactctatc	acggttaccc	agaacctaa	aagatgagtg	ttttgctaa	aaccaagaat	550
ctcaactatg	agtatgatgg	tattatgcag	aaatctcaag	ataatgtcac	agaactgtac	600
gaggttttca	tcagcttgct	tgtctcattc	cctgatgtta	cgagcaatat	gacatcttc	650
tgatattctg	aaaactgaca	gaagcgctgt	ttatcttcac	ctttctctat	agagcttgag	700
gacccctcgc	ctcccccaga	ccacattcct	tggattacag	ctgtacttcc	aacagttatt	750
atattctcga	tgggtttctg	tctaattcta	tggaaatgga	agaagaagaa	ggggcctcgc	800
gactcttata	aatgttggaac	caacacaaatg	gagaggggaa	agagtgaaca	gaccaagaaa	850
agagaaduaa	tcacataacc	tgaaagatct	gatgaagccc	agcgtgtttt	taaaagtctg	900
agagacattt	catgcgacaa	aagtgataca	tgtttttaat	taaagagtaa	agcccataca	950
agtatttcatt	ttttctaccc	tttcttttgt	aagttcctgg	gcaacctttt	tgatttcttc	1000
cagatggcnaa	aaagacatta	ccatgagtaa	taagggggct	ccaggactcc	ctctaagtgg	1050
aataccctcc	ctgtaactcc	agctctgctc	cgtatgccaa	gaggagactt	taattctctt	1100
actgcttctt	ttcaactcag	agcacactta	tgggcacaagc	ccagcttaat	ggctcatgac	1150
ctggcaaaaa	aatttaggac	caataaaaaaa	aaaaaaaaaa	aaaa		1200

SI100: 27

SI110: 213

SI120: PHT

SI130: Homo sapiens

M400-22

Met Gly Leu Ser Asn Ile Leu Phe Val Met Ala Phe Leu Leu Ser Gly  
 1 5 10 15  
 Ala Ala Pro Leu Lys Ile Gln Ala Tyr Phe Asn Glu Thr Ala Asp Leu  
 20 25 30  
 Pro Cys Gln Phe Ala Asn Ser Gln Asn Gln Ser Leu Ser Glu Leu Val  
 35 40 45  
 Val Phe Trp Gln Asp Gln Glu Asn Leu Val Leu Asn Glu Val Tyr Leu  
 50 55 60  
 Gly Lys Glu Lys Phe Asp Ser Val His Ser Lys Tyr Met Gly Arg Thr  
 65 70 75 80  
 Ser Phe Asp Ser Asp Ser Trp Thr Leu Arg Leu His Asn Leu Gln Ile  
 85 90 95  
 Lys Asp Lys Gly Leu Tyr Gln Cys Ile Ile His His Lys Lys Pro Thr  
 100 105 110  
 Gly Met Ile Arg Ile His Gln Met Asn Ser Glu Leu Ser Val Leu Ala  
 115 120 125  
 Asn Phe Ser Gln Pro Glu Ile Val Pro Ile Ser Asn Ile Thr Glu Asn  
 130 135 140  
 Val Tyr Ile Asn Leu Thr Cys Ser Ser Ile His Gly Tyr Pro Glu Pro  
 145 150 155 160  
 Lys Lys Met Ser Val Leu Leu Arg Thr Lys Asn Ser Thr Ile Glu Tyr  
 165 170 175  
 Asp Gly Ile Met Gln Lys Ser Gln Asp Asn Val Thr Glu Leu Tyr Asp  
 180 185 190  
 Val Ser Ile Ser Leu Ser Val Ser Phe Pro Asp Val Thr Ser Asn Met  
 195 200 205  
 Thr Ile Phe Cys Ile Leu Glu Thr Asp Lys Thr Arg Leu Leu Ser Ser  
 210 215 220  
 Pro Phe Ser Ile Glu Leu Glu Asp Pro Gln Pro Pro Pro Asp His Ile  
 225 230 235 240  
 Pro Trp Ile Thr Ala Val Leu Pro Thr Val Ile Ile Cys Val Met Val  
 245 250 255  
 Phe Lys Leu Ile Leu Trp Lys Trp Lys Lys Lys Arg Pro Arg Asn  
 260 265 270  
 Ser Tyr Lys Cys Gly Thr Asn Thr Met Glu Arg Glu Glu Ser Glu Gln  
 275 280 285  
 Thr Lys Lys Arg Glu Lys Ile His Ile Pro Glu Arg Ser Asp Glu Ala  
 290 295 300  
 Gln Arg Val Phe Lys Ser Ser Lys Thr Ser Ser Cys Asp Lys Ser Asp  
 305 310 315 320  
 Thr Cys Phe

M10-13

M11-1183

M12-DNA

M13-Mus musculus

M400-23

ggagcaagca gacgcgtaag agtggctcct gtaggcagca cggacttgaa caaccagact	60
actgtagacg tgttccagaa cttacggaag caccacagat ggaccccaga tgcaccatgg	120
gcttggaat ccttatcttt gtgacagtct tgcgtatctc agatgctgtt tccgtggaga	180
cgcaagotta tttaaatggg actgcatatc tgcggtgcc atttacaag gctcaaaaca	240

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taagcctgag tgagctggta gtattttggc aggaccagca aaagttgggt ctgtacgagc 300
actattttggg cacagagaaa cttgatagtg tgaatgpcaa gtacctgggc cgcacgagct 360
ttgacaggaa caactggact ctacgacttc acaatgttca gatcaaggac atgggctcgt 400
atgattgttt tatabaaaaa aagccaccca caggatcaat tatcctccaa cagacattaa 460
cagaactgtc agtgatcgcc aacttcagtg aactgaaat aaaaatggct cagaatgtaa 520
caggaaaattc tggcataaatt ttgacctgca cgtctaagca aggtcacccg aaacctaaag 580
agatgtattt tctgataact aattcaacta atgagtatgg tgataacatg cagatatcac 640
aagataaagt cacagaaactg ttcagtatct ccaacagcct ctctcttcca ttccgggatg 700
gtgtgtggga tatgacctt gtgtgtgttc tggaaaagg gtaaatgaag atttctccaa 760
aaactctcaa tttactcaa gattttccat ctctccaaac gtattggaag gagattacag 820
cttcagttaa tggggccctc ctcttggga tctgtctcat cattgtatgt cacaagaagc 880
cgaatcagcc tagcaggccc agaacacag cctctaagtt agagcgggat agtaacgctg 940
acagagagac tatcaacctg aaggaaactg aaccccaaat tcttcagca aaaccaaatg 1000
cagagtgaag gcagtgaag cctgaggaaa gaggtaaaaa ttgctttgcc tgaataaaga 1060
agtgcagagt tctcagaat tcaaaaatgt tctcagctga ttggaattct acagttgaat 1120
aattaaagaa caaatacac aacagtgaat aaaaaaaaaa aaa 1180

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1210 - 24

1211 - 209

1212 - FRP

1213 - Mus musculus

1400 - 14

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Met Asp Pro Arg Cys Thr Met Gly Leu Ala Ile Leu Ile Phe Val Thr
1 10 15
Val Leu Leu Ile Ser Asp Ala Val Ser Val Glu Thr Gln Ala Tyr Phe
20 25 30
Asn Gly Thr Ala Tyr Leu Pro Cys Pro Phe Thr Lys Ala Gln Asn Ile
35 40 45
Ser Leu Ser Glu Leu Val Val Phe Trp Gln Asp Gln Gln Lys Leu Val
50 55 60
Leu Tyr Glu His Tyr Leu Gly Thr Glu Lys Leu Asp Ser Val Asn Ala
65 70 75 80
Lys Tyr Leu Gly Arg Thr Ser Phe Asp Arg Asn Asn Trp Thr Leu Arg
85 90 95
Leu His Asn Val Gln Ile Lys Asp Met Gly Ser Tyr Asp Cys Phe Ile
100 105 110
Gln Lys Lys Pro Pro Thr Gly Ser Ile Ile Leu Gln Gln Thr Leu Thr
115 120 125
Glu Leu Ser Val Ile Ala Asn Phe Ser Glu Pro Glu Ile Lys Leu Ala
130 135 140
Gln Asn Val Thr Gly Asn Ser Gly Ile Asn Leu Thr Cys Thr Ser Lys
145 150 155 160
Gln Gly His Pro Lys Pro Lys Lys Met Tyr Phe Leu Ile Thr Asn Ser
165 170 175
Thr Asn Glu Tyr Gly Asp Asn Met Gln Ile Ser Gln Asp Asn Val Thr
180 185 190
Glu Leu Phe Ser Ile Ser Asn Ser Leu Ser Leu Ser Phe Pro Asp Gly
195 200 205
Val Trp His Met Thr Val Val Cys Val Leu Glu Thr Glu Ser Met Lys
210 215 220
Ile Ser Ser Lys Pro Leu Asn Phe Thr Gln Glu Phe Pro Ser Pro Gln
225 230 235 240
Thr Tyr Trp Lys Glu Ile Thr Ala Ser Val Thr Val Ala Leu Leu Leu

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245                      250                      255  
 Val Met Leu Leu Ile Ile Val Cys His Lys Lys Pro Asn Gln Pro Ser  
 260                      265                      270  
 Arg Pro Ser Asn Thr Ala Ser Lys Leu Glu Arg Asp Ser Asn Ala Asp  
 275                      280                      285  
 Arg Glu Thr Ile Asn Leu Lys Glu Leu Glu Pro Gln Ile Ala Ser Ala  
 290                      295                      300  
 Lys Pro Asn Ala Gln  
 305

K110-15  
 K111-1112  
 K112- DNA  
 K113- Homo sapiens

K400-15  
 cacaggtga aagctttagt tctctgtgc tctaacaggg actagacag acacacggat 50  
 gagggtggtc atttcagat attaggtac agcagaagca gcaaaaatgg atccccagt 100  
 cactatggga ctgagtaaca ttctcttctt gatgggttc ctgtctctgt gtgtgtgtcc 150  
 tctgaagatt ctagcttatt tcaatgagc tgcagacccg ccattgcaat ttgcacactc 200  
 tcaaaaccaa agcttgagtg agcttagtct attttggcag gacaggaaa acttggttct 250  
 gaatgaggtt taattaggca aagagaaatt tgcagtggtt cattccaagt atatgggccc 300  
 cacaagtttt gattcggaca gtgggaacct gagaactcac aatcttcaga tcaaggacaa 350  
 gggctgttat caatgttcca tccatcacia aaagcccaca ggaatgatto gcattccaca 400  
 gatgaattct gaactgtcag tctttgttaa ctccagtcac cctgaaatag taccatttcc 450  
 taatataaca gaaaatggtt acataaaatt gacctgtcca tctatacacg gtcaccaga 500  
 aactaaagag atgagtggtt tctaagaac caagaattca actatcgagt atgatggtat 550  
 tatgcacaaa tctcaagata atgtcacaga actgtaacgac gtttcaccca gcttgtctgt 600  
 ttcattctct gatgttaaga gcaatatgag catcttctgt attctggaaa ctgacaagac 650  
 gggcttitta tcttcacctt tctctataga gcttgaggac cctcagcttc cccagaccca 700  
 ccttccttgg attacagctg taattccaac agtctattata tctgtgatgg tttctgtct 750  
 aattctatgg aaatggaaga agagaaagcg gcttcgcaac tcttataaat gtcgaaccaa 800  
 caccatppag agggaaagaga gtgaacagac caagaaaaaga gaaaaaatcc atatactga 850  
 agatctgat gaagccacgc gtgtttttaa aagttogaag acatcttcat ggcacaaaag 900  
 ccaatcagtg ttttaattaa agtgtaaaag cc 1112

K110-16  
 K111-109  
 K112- FET  
 K113- Homo sapiens

K400-16  
 Met Asp Pro Gln Cys Thr Met Gly Leu Ser Asn Ile Leu Phe Val Met  
 1                      5                      10                      15  
 Ala Phe Leu Leu Ser Gly Ala Ala Pro Leu Lys Ile Gln Ala Tyr Phe  
 20                      25                      30  
 Asn Glu Thr Ala Asp Leu Pro Cys Gln Phe Ala Asn Ser Gln Asn Gln  
 35                      40                      45  
 Ser Leu Ser Glu Leu Val Val Phe Trp Gln Asp Gln Glu Asn Leu Val  
 50                      55                      60  
 Leu Asn Glu Val Tyr Leu Gly Lys Glu Lys Phe Asp Ser Val His Ser  
 65                      70                      75                      80  
 Lys Tyr Met Gly Arg Thr Ser Phe Asp Ser Asp Ser Trp Thr Leu Arg  
 85                      90                      95

Leu His Asn Leu Gln Ile Lys Asp Lys Gly Leu Tyr Gln Cys Ile Ile  
 100 105 110  
 His His Lys Lys Pro Thr Gly Met Ile Arg Ile His Gln Met Asn Ser  
 115 120 125  
 Glu Leu Ser Val Leu Ala Asn Phe Ser Gln Pro Glu Ile Val Pro Ile  
 130 135 140  
 Ser Asn Ile Thr Glu Asn Val Tyr Ile Asn Leu Thr Cys Ser Ser Ile  
 145 150 155 160  
 His Gly Tyr Pro Glu Pro Lys Lys Met Ser Val Leu Leu Arg Thr Lys  
 165 170 175  
 Asn Ser Thr Ile Glu Tyr Asp Gly Ile Met Gln Lys Ser Gln Asp Asn  
 180 185 190  
 Val Thr Glu Leu Tyr Asp Val Ser Ile Ser Leu Ser Val Ser Phe Pro  
 195 200 205  
 Asp Val Thr Ser Asn Met Thr Ile Phe Cys Ile Leu Glu Thr Asp Lys  
 210 215 220  
 Thr Arg Leu Leu Ser Ser Pro Phe Ser Ile Glu Leu Glu Asp Pro Gln  
 225 230 235 240  
 Pro Pro Pro Asp His Ile Pro Trp Ile Thr Ala Val Leu Pro Thr Val  
 245 250 255  
 Ile Ile Cys Val Met Val Phe Cys Leu Ile Leu Trp Lys Trp Lys Lys  
 260 265 270  
 Lys Lys Arg Pro Arg Asn Ser Tyr Lys Cys Gly Thr Asn Thr Met Glu  
 275 280 285  
 Arg Glu Glu Ser Glu Gln Thr Lys Lys Arg Glu Lys Ile His Ile Pro  
 290 295 300  
 Glu Arg Ser Asp Glu Ala Gln Arg Val Phe Lys Ser Ser Lys Thr Ser  
 305 310 315 320  
 Ser Cys Asp Lys Ser Asp Thr Cys Phe  
 325